

Abstract

Methods to identify genes, the expression of which are altered in response to pathogen infection, are provided, as well as the genes identified thereby and their corresponding promoters.

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In re Patent Application of: Sheng Quan et al.

Title: PLANT GENES INVOLVED IN DEFENSE AGAINST PATHOGENS

Docket No.: 1360.003US2

Express Mail No. EL671641675US

Date Mailed: June 23, 2001